

XP-00144843

Q9y4b8.Trembl

P.D.	1-11-1999
P.	1

!!AA\_SEQUENCE 1.0

ID Q9Y4B8 PRELIMINARY; PRT; 468 AA.  
AC Q9Y4B8;  
DT 01-NOV-1999 (TrEMBLrel. 12, Created)  
DT 01-NOV-1999 (TrEMBLrel. 12, Last sequence update)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last annotation update)  
DE KIAA0772 PROTEIN.  
GN KIAA0772.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
RN [1]  
RP SEQUENCE FROM N.A.  
RC TISSUE=BRAIN;

RX MEDLINE; 99087487.

RA Nagase T., Ishikawa K., Suyama M., Kikuno R., Miyajima N., Tanaka A.,  
RA Kotani H., Nomura N., Ohara O.;  
RT "Prediction of the coding sequences of unidentified human genes. XI.  
RT The complete sequences of 100 new cDNA clones from brain which code  
RT for large proteins in vitro.";  
RL DNA Res. 5:277-286(1998).  
OR EMBL; AB018315; BAA34492.1; -.  
DR INTERPRO; IPR000648; -.  
DR PFAM; PF01237; Oxysterol\_BP; 1.  
DR PROSITE; PS01013; OSBP; 1.  
SQ SEQUENCE 468 AA; 53971 MW; 9C51392E0EDF4DF8 CRC64;

Q9y4b8 Length: 468 August 11, 19100 11:16 Type: P Check: 5128 ..

1 MNGEEEFFDA VTEANQKVTG MIDLDTSKNN RIGKTGERPS QENGIQKHRT

51 SLPAPMFSRS DFSVWTILKK CVGLELSKIT MPIAFNEPLS FLQRITEYME  
101 HVYLIHRASC QPQPLERMQS VAAFAVSAVA SQWERTGKPF NPLLGETYEL  
151 IREDLGFRFI SEQVSHHPPPI SAFHSEGLNH DFLFHGSIYP KLKFWGKSVE  
201 AEPRGTITLE LLKHNEAYTW TNPTCCVHNV IIGKLWIEQY GTVEILNHRT  
251 GHKCVLHFKP CGLFGKELHK VEGHIQDKNK KKLFMIYGKW TECLWGIDPV  
301 SYESFKKQER RGDHRLRKAKL DEDSGKADSD VADDVPVAQE TVQVIPGSKL  
351 LWRINTRPPN SAQMYNFTSF TVSLNELETG MEKTLPPPTDC RLRPDIRGME  
401 NGNMDLASQE KERLEEKQRE ARRERAKEEA EWQTRWFYPG NNPYTGTPDW  
451 LYAGDYFERN FSDCPDIY